## SEQUENCE LISTING

<110> Singh, Yogendra Khanna, Hemant

<120> A NOVEL PROTEIN MOLECULE USEFUL FOR INHIBITION OF ANTHRAX TOXIN

<130> 4752 103.1 US

<150> 09/821,348

<151> 2001-03-29

<160> 6

<170> FastSEQ for Windows Version 3.0

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<211> 735

<212> PRT

<213> Artificial Sequence

<220>

<223> Derived from B. anthracis.

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Lys Trp Ser Thr Ala Ser Asp Pro Tyr Ser Asp Phe Glu Lys Val Thr

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230
                                         235
Gly Arg Ile Asp Lys Asn Val Ser Pro Glu Ala Arg His Pro Leu Val
                245
                                     250
Ala Ala Tyr Pro Ile Val His Val Asp Met Glu Asn Ile Ile Leu Ser
                                 265
Lys Asn Glu Asp Gln Ser Thr Gln Asn Thr Asp Ser Gln Thr Arg Thr
                             280
                                                 285
Ile Ser Lys Asn Thr Ser Thr Ser Arg Asp Ala Asn Thr Val Gly Val
                        295
Ser Ile Ser Ala Gly Tyr Gln Asn Gly Phe Thr Gly Asn Ile Thr Thr
                    310
                                         315
Ser Ala Gly Phe Ser Asn Ser Asn Ser Ser Thr Val Ala Ile Asp His
                325
                                     330
Ser Leu Ser Leu Ala Gly Glu Arg Thr Trp Ala Glu Thr Met Gly Leu
                                 345
Asn Thr Ala Asp Thr Ala Arg Leu Asn Ala Asn Ile Arg Tyr Val Asn
                            360
Thr Gly Thr Ala Pro Ile Tyr Asn Val Leu Pro Thr Thr Ser Leu Val
                        375
                                            380
Leu Gly Lys Asn Gln Thr Leu Ala Thr Ile Lys Ala Lys Glu Asn Gln
                    390
                                        395
Leu Ser Gln Ile Leu Ala Pro Asn Asn Tyr Tyr Pro Ser Lys Asn Leu
                                    410
Ala Pro Ile Ala Leu Asn Ala Gln Asp Asp Phe Ser Ser Thr Pro Ile
                                425
Thr Met Asn Tyr Asn Gln Phe Leu Glu Leu Glu Lys Thr Lys Gln Leu
                            440
Arg Leu Asp Thr Asp Gln Val Tyr Gly Asn Ile Ala Thr Tyr Asn Phe
                        455
Glu Asn Gly Arg Val Arg Val Asp Thr Gly Ser Asn Trp Ser Glu Val
                    470
                                        475
Leu Pro Gln Ile Gln Glu Thr Thr Ala Arg Ile Ile Phe Asn Gly Lys
                485
                                    490
Asp Leu Asn Leu Val Glu Arg Arg Ile Ala Ala Val Asn Pro Ser Asp
            500
                                505
Pro Leu Glu Thr Thr Lys Pro Asp Met Thr Leu Lys Glu Ala Leu Lys
Ile Ala Phe Gly Phe Asn Glu Pro Asn Gly Asn Leu Gln Tyr Gln Gly
                        535
Lys Asp Ile Thr Glu Phe Asp Phe Asn Phe Asp Gln Gln Thr Ser Gln
                    550
                                        555
Asn Ile Lys Asn Gln Leu Ala Glu Leu Asn Ala Thr Asn Ile Tyr Thr
                565
                                    570
Val Leu Asp Lys Ile Lys Leu Asn Ala Lys Met Asn Ile Leu Ile Arg
                                585
Asp Lys Arg Phe His Tyr Asp Arg Asn Asn Ile Ala Val Gly Ala Asp
                            600
Glu Ser Val Val Lys Glu Ala His Arg Glu Val Ile Asn Ser Ser Thr
                        615
Glu Gly Leu Leu Asn Ile Asp Lys Asp Ile Arg Lys Ile Leu Ser
                    630
                                        635
Gly Tyr Ile Val Glu Ile Glu Asp Thr Glu Gly Leu Lys Glu Val Ile
                645
                                    650
Asn Asp Arg Tyr Asp Met Leu Asn Ile Ser Ser Leu Arg Gln Asp Gly
                               665
Lys Thr Phe Ile Asp Phe Lys Lys Tyr Asn Asp Lys Leu Pro Leu Tyr
        675
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Ile Ser Asn Pro Asn Tyr Lys Val Asn Val Tyr Ala Val Thr Lys Glu
                         695
                                             700
Asn Thr Ile Ile Asn Pro Ser Glu Asn Gly Asp Thr Ser Thr Asn Gly
                     710
                                         715
Ile Lys Lys Ile Leu Ile Phe Ser Lys Lys Gly Tyr Glu Ile Gly
                                     730
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      <223> Derived from B. anthracis.
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Phe Thr Gly Asn Ile Thr Thr
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      <210> 3
      <211> 23
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Thr His Thr Ser Glu Val His Gly Asn Ala Glu Val His Ala Ser Phe
                                     10
Phe Asp Ile Gly Gly Ser Val
            20
      <210> 4
      <211> 69
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atcactaca
                                                                      69
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      <213> Artificial Sequence
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      <223> Synthetic.
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attactaaat cctgcagatg tagtgatatt accagtaaag ccgttctgat accctgctga
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aattgaaact cctacagtat tagcatccct acttgtagaa gtatttttac
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<211> 23 <212> DNA

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23